Fig. 1A

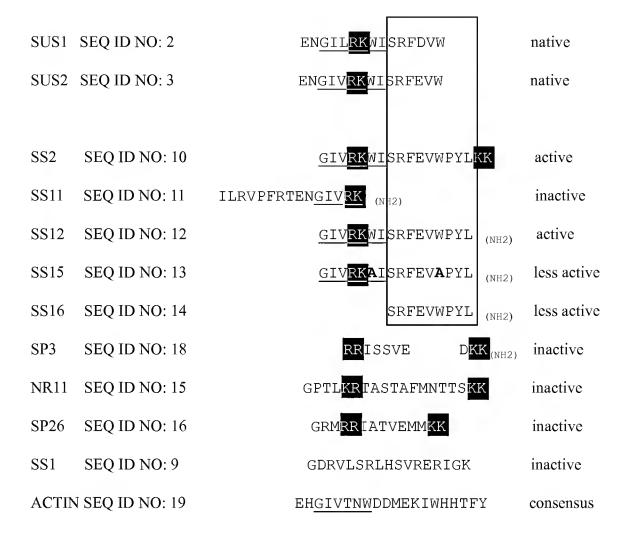|			
SUS1	SEQ ID NO: 2	ENGIL**RK**WISRFDVW	native
SUS2	SEQ ID NO: 3	ENGIV**RK**WISRFEVW	native
SS2	SEQ ID NO: 10	GIV**RK**WISRFEVWPYL**KK**	active
SS11	SEQ ID NO: 11	ILRVPFRTENGIV**RK** (NH2)	inactive
SS12	SEQ ID NO: 12	GIV**RK**WISRFEVWPYL (NH2)	active
SS15	SEQ ID NO: 13	GIV**RK**AISRFEV**A**PYL (NH2)	less active
SS16	SEQ ID NO: 14	SRFEVWPYL (NH2)	less active
SP3	SEQ ID NO: 18	**RR**ISSVE D**KK**(NH2)	inactive
NR11	SEQ ID NO: 15	GPTL**KR**TASTAFMNTTS**KK**	inactive
SP26	SEQ ID NO: 16	GRM**RR**IATVEMM**KK**	inactive
SS1	SEQ ID NO: 9	GDRVLSRLHSVRERIGK	inactive
ACTIN	SEQ ID NO: 19	EHGIVTNWDDMEKIWHHTFY	consensus

Double basic cluster: black box; e.g. **KK**
Possible region of specificity: underlined or boxed
Substitutions: bold